

1 AGATGAGTGT GGGGGTGA AGAGTCAAGT TCGGGGCAT  
 METSerVal GlyArgArg ArgValLysLeu LeuGlyIle  
 41 CCGATGATG GCAAAGTCT TCATTATTT GATTGTGAA  
 eLeuMETMET AlaAsnValPhe IleTyrLeu IleValGlu  
 81 GTCTCCAAA ACAGTAGCA AGACAAAAT CGAAGGGAG  
 ValSerLysAsn SerSerGln AspLysAsn GlyLysGlyG  
 121 GGTATATAT CCGAAGAG AGTTCCTGA AGCAACCCG  
 lyValIleIle ProLysGlu LysPheTrpLys ProProSe  
 161 CAGTCCCGG GGTACTGCA ACAGGGACA GGAAGCTG  
 rThrProArg AlaTyrTrpAsn ArgGluGln GluLysLeu  
 201 AACAGTGGT ACAATCCCT CTGACAGG GTGGCAATC  
 AsnArgTrpTyr AsnProIle LeuAsnArg ValAlaAsnG  
 241 AGACAGGGA GGTAGGACA TGTCCAAACA CAGTCACT  
 lThrGlyGlu LeuAlaThr SerProAsnThr SerHisLe  
 281 CAGCATTTG GAGCAGACT CGAGGTGAT GACAGCTG  
 uSerTyrCys GluProAspSer ThrValMET ThrAlaVal  
 321 ACAGATTTA ATAATCTGC GACAGATT AAAGACTTC  
 ThrAspPheAsn AsnLeuPro AspArgPhe LysAspPheL  
 361 TCTGTATTT GAGTGGCG AATTACTGC TCTTATAGA  
 euLeuTyrLeu ArgCysArg AsnTyrSerLeu LeuIleAs  
 401 TCAACCGAG AATGTGCAA AGAGCCCTT CTACTATTG  
 pGlnProLys LysCysAlaLys LysProPhe LeuLeuLeu  
 441 GCGATAAAGT CCGTATTC ACATTTGCC AGAAGGCGAG  
 AlaIleLysSer LeuIlePro HisPheAla ArgArgGlnA  
 481 CAATCCGGA GTCTGGGGC CGAGAAACA ACGTAGGGA  
 laIleArgGlu SerTrpGly ArgGluThrAsn ValGlyAs  
 521 CCGACAGTA GTGAGGGCT TCGGTGCG CAGACACC  
 nGlnThrVal ValArgValPhe LeuLeuGly LysThrPro  
 561 CCGAGGACA ACCACCTCA CTTTGGAC ATGCTAAGT  
 ProGluAspAsn HisProAsp LeuSerAsp METLeuLysP  
 601 TTGAGGTGA CAGCACCG GACATCTCA TGTGGACTA  
 heGluSerAsp LysHisGln AspIleLeuMET TrpAsnTy  
 641 TAGAGACACA TCTTCAACC TGACCTGAA GAGAGTCTG  
 rArgAspThr PhePheAsnLeu SerLeuLys GluValLeu

FIG. 1 (sheet 1 of 2)

641 TTCTCAGGT GGGTGGGAC TTCTGTCCA GAGCAGGT  
 PheLeuArgTrp ValSerThr SerDysPro AspAlaGluP  
 721 TTGTCTCAA GGGCATCAT GAGGTTTG TGAACACCA  
 heValPheLys GlyAspAsp AspValPheVal AsnThrHi  
 761 TCACATCCTT AATTACTTCA ATAGCTTATC CAAGACCAA  
 sHisIleLeu AsnTyrLeuAsn SerLeuSer LysSerLys  
 801 GCGAAGACT TGTTCATAGG TGGGTGATC CACAATGCTG  
 AlaLysAspLeu PheIleGly AspValIle HisAsnAlaG  
 841 GGGCTCAGG GGTATAGAA CTCAGTACT ACATCCAGA  
 LyProHisArg AspLysLys LeuLysTyrTyr IleProGl  
 881 AGTCTCTTC ACCGGGCTT ACCACCTTA TCGGGGGCT  
 uValPheTyr ThrGlyValTyr ProProTyr AlaGlyGly  
 921 GGTGGTTCC TGTACCCCG CCGCTTCC TTGAGGCTT  
 GlyGlyPheLeu TyrSerGly ProLeuAla LeuArgLeuT  
 961 ACAGTCCGC TGGGGGGTC CTCCTCTCC CTATTGATCA  
 yrSerAlaThr SerArgVal HisLeuTyrPro IleAspAs  
 1001 TGTCTATAG CCAATGTGCC TTGAGAACT GGGCTTGT  
 pValTyrThr GlyMetCysLeu GlnLysLeu GlyLeuVal  
 1041 CCGAGGAGC ACAAGGCTT CAGGATTT GATATTGAG  
 ProGluLysHis LysGlyPhe ArgThrPhe AspIleGluG  
 1081 AGAAAAATAA GAAAAATATT TGTCTCTATA TAGCCTAAT  
 LuLysAsnLys LysAsnIle CysSerTyrIle AspLeuME  
 1121 GTTAGTACAT AGCAGAAAC CTCAGAGAT GATTGATATC  
 TLeuValHis SerArgLysPro GlnGluMET IleAspIle  
 1161 TGTCTCTGT TCGAAGTCC TAATTAAAA TGTCTA  
 TrpSerGlnLeu GlnSerPro AsnLeuLys Cys

FIG. 1 (sheet 2 of 2)

```

MAX--PQKV-----LLR-LL--V
      10              20              30
MSVGP-RRV-----K-L
MA--S-SC-----Y-
MAP-----AVLTALPNRMSLPS-KWSL
MQSKHRL-----LFCIL--V
MLQWPRRHCCFAKMTWNAKRSIFPTHIGV

LSLVKLLXXXFXFLKH--W-----
      40              50              60
LGILMMANV-IY-IVEVSKNSSQDKNGKGG
LSVVC-----ASA-----
L-----LSSLSSLV-----
LPLILLVVGYCGLITHLH-----
LSIVFFIFAMELEFN-HD-LPGRAGFKENPV

--F--F--D-----TWY
      70              80              90
VIIPKEKWK-PSTPRAYWNREQEKLNRN
-----LWY
ELNFERHSHY-----
TYTERG-ERSTKSETNHSSLR-----NKK
LSIP--LRPQTGSXSXSXXLSHL-Y---N
      100             110             120
NPILNRVANGT-ELATSPNTSHLSVCEPDS
LSIT--P-TS-YTG-KPFSHI-----
LSLPHY-----
-----ENDDTGGGA-SGLDKFAV-----
ETVFOQLPPOTATNENNTOLSPQGVGTLEN
TVXRYVXXFNXXXT------PINSXXFEF
      130             140             150
TVMTAVTDFFNNLPDFFKDFLLYLRCRNYSL
TVARKNFTFGNIR------PHSPHF
VIEPVNWMYF-VEYE-----YRQDFRF
-----LRVPEFTAHV
TLSANGSIY-EKGIG-----HPNHYH-KY
LIDEPYKCKKK-PFLVLLIKSXPGXFXARQ
      160             170             180
LIFQFKKCAK-PPFLAIP-LIPHARRC
LINEFNCEFNIPFLI-ISTTHKE-DARC
TLRFHSN-SHQNEFLITVT-RSDVKARC
PVGQEAR-----TMEIAVGNRRPE
LINEFEKQEQE-SPFI-IAAEEQIEER
AIRETWGXEXNFXGIXVXRVELLGKXA-EX
      190             200             210
AIRESWGRETNVGNOTVVEVFLLGFTTPFD
AIRETWGDFNNHKGK-KIATLFLLGNN-
AIPVTWGEKKSWWGYEMLTFELGQQA-FR
AIFRTWGYEGRESDVHLRVELLGTAEDE
AIFOTWGNFSLAPG-QIT-IFELLSI--K
XDPXLXXMVEXESRXHGDIIQQDFLDITYFN
      220             230             240
NHEDLSDMLKFEESDKHQDILMWNRYD-FR
-LFVLNQMVQESQIFHDLIVEDFISYHN
E-KTIALSLFDEHVLVJDI-RDDFISTYNN
K------WAWESHEDILCADSTCAYEN
LNGYIQRAILEESQYHDTIOGEYDITYY
LTLKTLMGMRWVATFCPXA EYVMKTDSDVF
      250             260             270
SLKEVFLFLRWASISCHDAFFFGGLDDCV
TLKKTTEGMRWVATFCISKAKYVWAVD-LIF
TLKKTIFAFRWMEFCINAKVIMKTTDVE
NTLKTMLGHPWASEQFNRSFYLFVLDEYY
IKKTEMGMNWVATYCHIFVWVWDDSTME
VNTXNLLNKKLLKPSLSHRXXLFTGY-VIXG
      280             290             300
VNTHHIINHYLNSLKKKAKKDFIG-DVINH
VHMONLIYEL-ESTKPRRYFTGY-VIN
IN-GNIVKYLI--N-NHSEKFFFTVPLTDN
VSAKNVVKELGRGROFQOPELFAGHAFQT
VNTHEYLINELI-DEPPHNHYFTGY-LMR

```

FIG. 2 (sheet 1 of 2)

YGPYDCKFSKWTLPKDLYPFYVYPPYCSGG  
 310 320 330  
 AGIHRDYLKLYYFVEVLT-GYPPYAGG  
 -GIRIVRSGKMYMERLYYDSNYPPFCST  
 S-YPGFHHKNNHISYQEVPIKFFPPYCCCL  
 -SHLFHKEKSWYVSLLEYPFDRWPPVTA  
 VAFNENFDSKSWYMIPTLYFSERVEVF  
 T  
 GYIFSGDLAERLYKASLHVRLHLEDVYVG  
 340 350 360  
 FLYCPIALHINSTSRHLYPID  
 VHSALVALIY-T  
 THGDDVPVYEMMSKPIKF  
 AFGLSQKALRQAAVLPFRFDVYL  
 VEGHDAKIFVGI  
 ICLXKLGIDPXXPXG--FNHW-KXXKSXC  
 370 380 390  
 MCLQKLGGLVSEKHKCFRTSDISEKKNKNI  
 LCLRKLGHFQNSQ--FHHF--FMAYLC  
 -YNLFKVEIHICEDTNLFPLY-RIHLDV  
 VALHAGISLOHCDD--RFHRPAYGPD  
 TIAKIRTDVPEPNEFV--RVSYST  
 SYSRVIAV4OF-SPEEMIRIWNXL-Q-KNL  
 400 410 420  
 YIDLMLVSR-KQOEN-DILSOI-SP  
 RYRRVITVHAI-UPHEHPIHDMSSKH  
 QLRPVIAHGF-SKEITFMQVML--RT  
 YSSVITASHFEGDPEETVWE--  
 KYCHLITSHGF-QISELTKYHMHQNH  
 XC-----Y-----  
 430  
 KC  
 RC  
 TC-----H  
 -C-----RSANY-----A  
 ACANAAKEXAGRVRHRKLH

FIG. 2 (sheet 2 of 2)

000435 300000

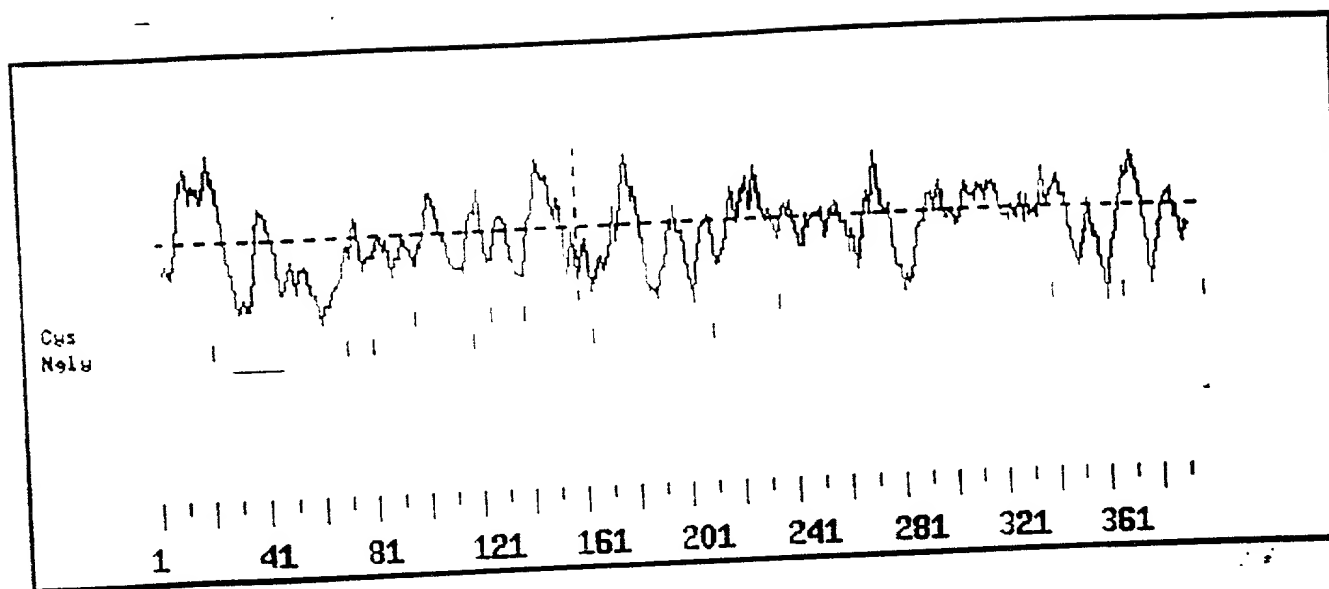


FIG. 3



581 CCCCCCTCCG ATGGACAAGC TCATAAGTGA CCCCCCTATC  
 eProValLeu METAspLysLeu IleSerAsp LeuProIle  
 721 TCCAGGAGAC ACAATTTTAT GCCTCCCTTA CCCAATATCA  
 TyrArgArgHis AsnPheMET ValSerLeu ProAsnIleT  
 761 CAGATTCAGT CATGAAAAG AAGGCGCAAT TTCGCAAGCA  
 hrAspSerVal IleGluLys LysArgGlnPhe LeuLysGI  
 801 RAGGATTTCG CTGGAGCGAT TTCCTGCTCA CCTAGTGAAT  
 nArgIleTrp LeuGluGlyPhe AlaAlaAsp LeuValAsn  
 841 ATCATCCCTT CTCGCGCTT TCCCTGGAC AGTGATTTCG  
 IleIleProSer LeuIhrPhe LeuLeuAsp SerAspLeuG  
 881 AGCTCTCTAA GAAAGCATG AATTCCTAC GCCTGTGTT  
 LuIhrLeuLys LysSerMET LysPheTyrArg ThrValPh  
 921 TCGAGTCCAT CAAACTCTT TCGGAGATT AGCTAGGGAC  
 eGlyValAsp GluIhrSerLeu GlnArgLeu AlaArgAsp  
 961 TCGGAATAG AGGTGGTCA CGTGGAGCC ATGATAAAAT  
 TrpGluIleGlu ValAspGln ValGluAla METIleLysS  
 1001 CTCCTCCCTG GTTCAAACT ACAGATGAG AAACAATACA  
 erProAlaVal PheLysPro ThrAspGluGlu ThrIleGI  
 1041 AGAAGGCTT TCAAGATATA TTCAGGATT CTGTTCGCT  
 nGluArgLeu SerArgTyrIle GlnGluPhe CysLeuAla  
 1081 AATGGTACT TACTTCTAA AATAGTTTT CTAAAGAAA  
 AsnGlyTyrLeu LeuProLys AsnSerPhe LeuLysGluI  
 1121 TATTTTACT GAAATATAT TTCCTGACA TGGTCACTCA  
 lePheTyrLeu LysTyrTyr PheLeuAspMET ValThrGI  
 1161 CGATGCTAAA ACTCTCTTA AAGAGATAG TTAAAGAAC  
 AspAlaLys ThrLeuLeuLys GluIleCys LeuArgAsn  
 1201 PG

FIG. 4 (sheet 2 of 2)

Cus  
N91y

FIG. 5



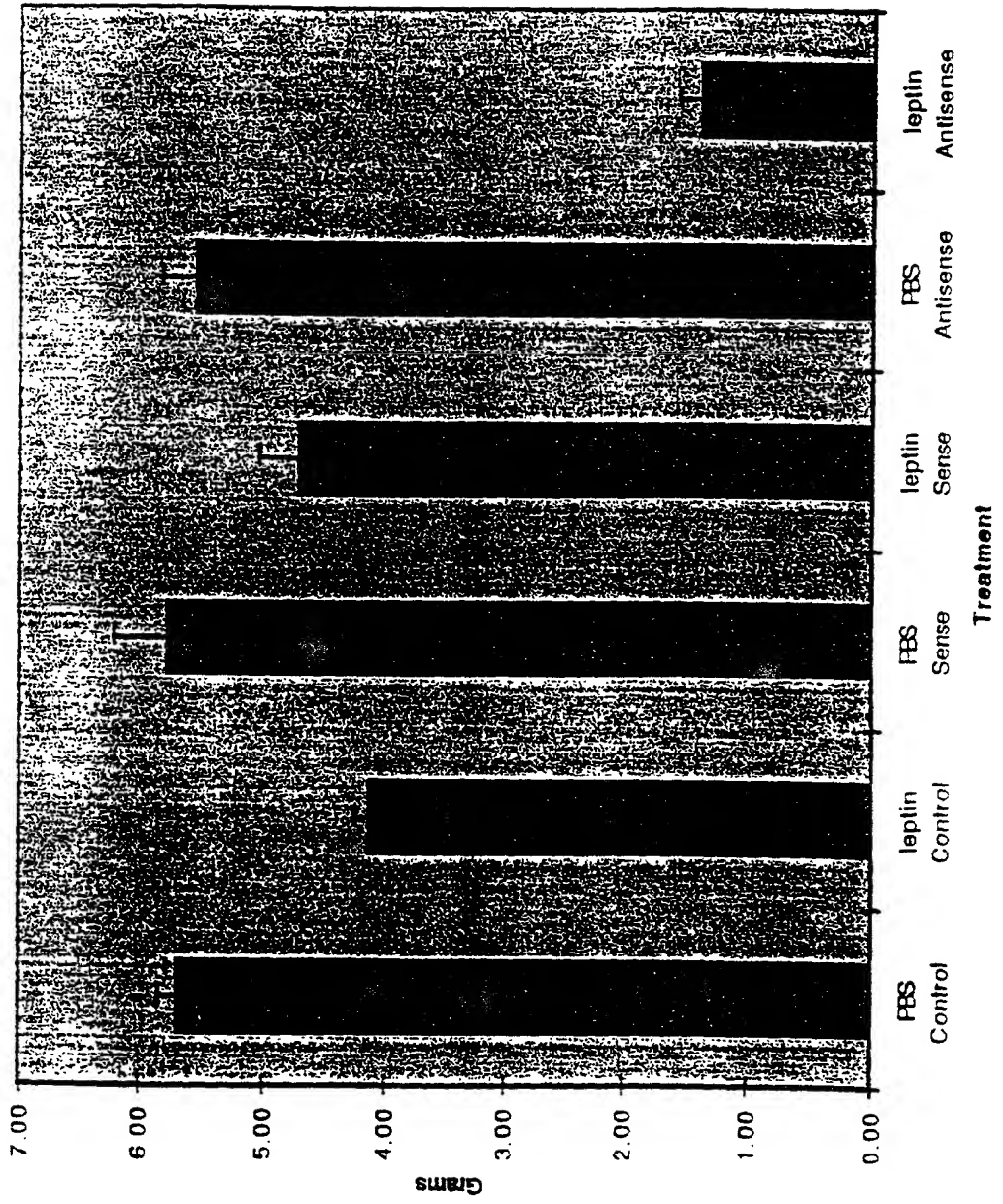


FIG. 6

5'

ACGCGTCCGCGCAGCGGCAGCGGCAGCAGCGGCAACAAGTGCCGGAGGCTA  
GCAGAGCCAAGCCGGAGCAGTCCCTGCCGCCGACACCGCCGGGGCCCGCCGTC  
CGGGGCGCCGCGCATGGAGCGTGAGCTGCCGCCGGTCGCCGGGCTGAGCCGC  
GCGGAGCGGCCGGGACGTGGATGTGGCCGCGATCTCCCGCCCTTGCCCCCGC  
CCCGCCGAGCTGGAGCTGCTCCCGGACAAGATATGAGAAAT**T**GAGTGTTGGA  
CGTCGAAGAATAAAGTTGTTGGGTATCCTGATGATGGCAAATGTCTTCATTTA  
TTTTATTATGGAAGTCTCCAAAAGCAGTAGCCAAGAAAAAAATGGAAAAGGG  
GAAGTAATAATACCCAAAGAGAAGTTCTGGAAGATATCTACCCCTCCCGAGG  
CATACTGGAACCGAGAGCAAGAGAAGCTGAACCGGCAGTACAACCCCATCCT  
GAGCATGCTGACCAACCAGACGGGGGAGGCGGGCAGGCTCTCCAATATAAG  
CCATCTGAACTACTGCGAACCTGACCTGAGGGTCACGTCGGTGGTTACGGGT  
TTTAACAACCTTGCCGGACAGATTTAAAGACTTTCTGCTGTATTTGAGATGCCG  
CAATTATTCAGTCTTATAGATCAGCCGGATAAGTGTGCAAAGAAACCTTTCT  
TGTTGCTGGCGATTAAGTCCCTCACTCCACATTTTGCCAGAAGGCAAGCAATC  
CGGGAATCCTGGGGCCAAGAAAGCAACGCAGGGAACCAAACGGTGGTGCGA  
GTCTTCCTGCTGGGCCAGACACCCCCAGAGGACAACCACCCCGACCTTTTCA  
ATATGCTGAAATTTGAGAGTGAGAAGCACCAAGACATTCTTATGTGGAAC  
CAGAGACACTTTCTTCAACTTGTCTCTGAAGGAAGTGCTGTTTCTCAGGTGGG  
TAAGTACTTCCTGCCAGACACTGAGTTTGTGTTTCAAGGGCGATGACGATGTT  
TTTGTGAACACCCATCACATCCTGAATTACTTGAATAGTTTATCCAAGACCAA  
AGCCAAAGATCTCTTCATAGGTGATGTGATCCACAATGCTGGACCTCATCGG  
GATAAGAAGCTGAAGTACTACATCCCAGAAGTTGTTTACTCTGGCCTCTACCC  
ACCCTATGCAGGGGGAGGGGGGTTTCTCTACTCCGGCCACCTGGCCCTGAGG  
CTGTACCATATCACTGACCAGGTCCATCTCTACCCCATTTGATGACGTTTATAC  
TGGAATGTGCCTTCAGAACTCGGCCTCGTTCCAGAGAAACACAAAGGCTTC

AGGACATTTGATATCGAGGAGAAAAACAAAAATAACATCTGCTCCTATGTAG  
ATCTGATGTTAGTACATAGTAGAAAACCTCAAGAGATGATTGATATTTGGTCT  
CAGTTGCAGAGTGCTCATTTAAAATGCT**T**AAAATAGATACAAACTCAATTTKG  
SATWGRAAGGGGTWTTTTGRATWGGYCCCATGTTGGGGTCTCACATTAGAGT  
AATTTCTATTTNAANCATGAAATTGCCTTTATGAGTGATACCCATTTANGGCC  
TCTAANCCTTCATTTGNACTCACGTGAAGAAGGGAAAGCGGGAGAAGGTAAT  
TTNTTTATGGTGAATGGCAGGATATTGGTCTGACTTACCGNTAGGGGANTTTA  
AAACTGGNCCTTTTTGAATCTGTTTGGATGGCCCTT

MSVGRRRIKLLGILMMANVFIYFIMEVSKSSSQEKNGKGEVIIPKEKFWKISTPPE  
AYWNREQEKLNRQYNPILSMLTNQTGEAGRLSNISHLNYCEPDLRVTSVVTGFN  
NLPDRFKDFLLYLRCRNYSLIDQPDKCAKKPFLLAIAKSLTPHFARRQAIRESWG  
QESNAGNQTVVVRVFLLGQTPPEDNHPDLSDMLKFESEKHQDILMWNRYRDTFFNL  
SLKEVLFLRWVSTSCPDTEFVFKGDDDVFNTHHILNYLNSLSKTKAKDLFIGDV  
IHNAGPHRDKKLKYYIPEVVYSGLYPPYAGGGGFLYSGHLALRLYHITDQVHLY  
PIDDVYTGMCLOKLGLVPEKHKGFRTFDIEEKNKNNICSYVDLMLVHSRKPQEM  
IDIWSQLQSAHLKC

MSVGRRRIKLLGILMMANVFIYFIMEVSKSSSQEKNGKGEVIIPKEKFWKISTPPE

START

201 CCCCCCCCCCGAGCTGGAGCTGCTCCCGGACAAGATATGAGAAATGAGTGTGGGACGT 260  
 5 CCCCCCCCCCGAGCTGGAGGTGTCCCTAGACAAGGTATGAGAGATGAGTGTGGGGCGT 64  
 351 CGAAGAATAAAGTTGTTGGGTATCCTGATGATGGCAAATGTCTTCATTTATTTTATTATG 320  
 65 CGAAGAGTCAAGTTGCTGGGCATCCTGATGATGGCAAATGTCTTCATTTATTTGATTGTG 124  
 321 GAAGTCTCCAAAAGCAGTAGCCAAGAAAAAATGGAAAAGGGGAAGTAATAATACCCAAA 380  
 125 GAAGTCTCCAAAACAGTAGCCAAGACAAAAATGGAAAAGGGAGGAGTAATAATCCCGAAA 184  
 381 GAGAAGTTCTGGAAGATATCTACCCCTCCCGAGGCATACTGGAACCGAGAGCAAGAGAAG 440  
 185 GAGAAGTTCTGGAAGCCACCCAGCACTCCCGGGCATACTGGAACAGGGAACAGGAGAAG 244  
 441 CTGAACCGGGCAGTACAACCCCATCCTGAGCATGCTGACCAACCAGACGGGGGAGGCGGGC 500  
 245 CTGAACAGGTGGTACAATCCCATCTTGAACAGGGTGGCCATCAGACAGGGGAGCTAGCC 304  
 501 AGGCTCTCCAATATAAGCCATCTGAACCTGCGAACCTGACCTGAGGGTCACGTGCGGTG 560  
 305 ACATCTCCAAACACAAGTCACCTGAGCTATTGTGAACCAAGACTCGACGGTCATGACAGCT 364  
 561 GTTACGGGTTTTTAACAACCTGCCGGACAGATTTAAAGACTTTCTGCTGTATTTGAGATGC 620  
 365 GTGACAGATTTTAATAATCTGCCGGACAGATTTAAAGACTTTCTCTGTATTTGAGATGC 424  
 621 CGCAATTATTCAGTCTTATAGATCAGCCGGATAAGTGTGCAAAGAAACCTTTCTTGTG 680  
 425 CGGAATTACTCGCTGCTTATAGATCAACCGAAGAAATGTGCAAAGAAGCCCTTCTTACTA 484  
 681 CTGGCGATTAAAGTCCCTCACTCCACATTTTGCCAGAAGGCAAGCAATCCGGGAATCCTGG 740  
 485 TTGGCGATAAAGTCCCTCATTCACATTTTGCCAGAAGGCAAGCAATCCGGGAGTCTTGG 544  
 741 GGCCAAAGAAAGCAACGCAGGGAACCAACGGTGGTGCGAGTCTTCCTGCTGGGCCAGACA 800  
 545 GGCCGAGAAACCAACGTAGGGAACCAGACAGTAGTGAGGGTCTTCCTGTTGGGCAAGACA 604  
 801 CCCCCAGAGGACAACCACCCGACCTTTTCAGATATGCTGAAATTTGAGAGTGAGAAGCAC 860  
 605 CCCCCAGAGGACAACCACCTGACCTTTTCGGACATGCTTAAGTTTGAGAGTGACAAGCAC 664  
 861 CAAGACATTCTTATGTGGAACCTACAGAGACACTTTCTTCAACTTGTCTCTGAAGGAAGTG 920  
 665 CAGGACATCCTCATGTGGAACCTATAGAGACACATTCTTCAACCTGTCCCTGAAGGAAGTG 724  
 921 CTGTTTCTCAGGTGGGTAAAGTACTTCCTGCCCAGACACTGAGTTTGTGTTTCAAGGGCGAT 980  
 725 CTGTTTCTTAGGTGGGTGAGCACTTCCTGTCCAGACGCGAGAGTTTGTCTTCAAGGGCGAT 784  
 981 GACGATGTTTTTGTGAACACCCATCACATCCTGAATTACTTGAATAGTTTATCCAAGACC 1040  
 785 GATGACGTGTTTGTGAACACCCATCACATCCTTAATTACTTGAATAGCTTATCCAAGACC 844  
 1041 AAAGCCAAAGATCTCTTCATAGGTGATGTGATCCACAATGCTGGACCTCATCGGGATAAG 1100  
 845 AAAGCCAAAGACTTGTTCATAGGTGACGTGATCCACAATGCTGGGCCTCACCGGGATAAG 904

FIG. 9 (10F2)



Variable	Mean	SD	Min	Max
Age	35.2	12.5	18	65
Gender	0.52	0.50	0	1
Marital status	0.68	0.48	0	1
Education	12.5	2.1	9	16
Income	15.2	8.5	5	35
Health status	0.75	0.43	0	1
Employment	0.82	0.38	0	1
Home ownership	0.71	0.45	0	1
Vehicle ownership	0.65	0.48	0	1
Life satisfaction	4.2	1.5	1	7
Health satisfaction	5.1	1.2	1	7
Income satisfaction	3.8	1.8	1	7
Home satisfaction	4.5	1.4	1	7
Vehicle satisfaction	4.0	1.6	1	7
Life satisfaction (cont.)	4.3	1.4	1	7
Health satisfaction (cont.)	5.2	1.1	1	7
Income satisfaction (cont.)	3.9	1.7	1	7
Home satisfaction (cont.)	4.6	1.3	1	7
Vehicle satisfaction (cont.)	4.1	1.5	1	7

Variable	Mean	SD	Min	Max
Age	35.2	12.5	18	65
Gender	0.52	0.50	0	1
Marital status	0.68	0.48	0	1
Education	12.5	2.1	9	16
Income	15.2	8.5	5	35
Health status	0.75	0.43	0	1
Employment	0.82	0.38	0	1
Home ownership	0.71	0.45	0	1
Vehicle ownership	0.65	0.48	0	1
Life satisfaction	4.2	1.5	1	7
Health satisfaction	5.1	1.2	1	7
Income satisfaction	3.8	1.8	1	7
Home satisfaction	4.5	1.4	1	7
Vehicle satisfaction	4.0	1.6	1	7
Life satisfaction (cont.)	4.3	1.4	1	7
Health satisfaction (cont.)	5.2	1.1	1	7
Income satisfaction (cont.)	3.9	1.7	1	7
Home satisfaction (cont.)	4.6	1.3	1	7
Vehicle satisfaction (cont.)	4.1	1.5	1	7

Variable	Mean	SD	Min	Max
Age	35.2	12.5	18	65
Gender	0.52	0.50	0	1
Marital status	0.68	0.48	0	1
Education	12.5	2.1	9	16
Income	15.2	8.5	5	35
Occupation	1.2	0.8	0	2
Health status	2.1	1.2	1	4
Stress level	3.5	1.5	1	5
Life satisfaction	4.2	1.8	1	7
Resilience	5.1	2.2	1	8
Optimism	6.3	2.5	1	9
Gratitude	7.2	2.8	1	10
Forgiveness	8.1	3.1	1	11
Empathy	9.0	3.4	1	12
Compassion	9.8	3.6	1	13
Kindness	10.5	3.8	1	14
Generosity	11.2	4.0	1	15
Patience	12.0	4.2	1	16
Self-control	12.8	4.4	1	17
Perseverance	13.5	4.6	1	18
Determination	14.2	4.8	1	19
Confidence	15.0	5.0	1	20
Trust	15.8	5.2	1	21
Cooperation	16.5	5.4	1	22
Teamwork	17.2	5.6	1	23
Leadership	18.0	5.8	1	24
Influence	18.8	6.0	1	25
Power	19.5	6.2	1	26
Authority	20.2	6.4	1	27
Control	21.0	6.6	1	28
Management	21.8	6.8	1	29
Organization	22.5	7.0	1	30
Planning	23.2	7.2	1	31
Execution	24.0	7.4	1	32
Monitoring	24.8	7.6	1	33
Evaluation	25.5	7.8	1	34
Reflection	26.2	8.0	1	35
Learning	27.0	8.2	1	36
Growth	27.8	8.4	1	37
Development	28.5	8.6	1	38
Progress	29.2	8.8	1	39
Success	30.0	9.0	1	40

Variable	Mean	SD	Min	Max
Age	35.2	12.5	18	65
Gender	0.52	0.50	0	1
Marital status	0.68	0.48	0	1
Education	12.5	2.1	9	16
Income	15.2	8.5	5	35
Health status	0.75	0.43	0	1
Employment	0.82	0.38	0	1
Home ownership	0.71	0.45	0	1
Vehicle ownership	0.65	0.48	0	1
Life satisfaction	4.2	1.5	1	7
Health satisfaction	5.1	1.2	1	7
Income satisfaction	3.8	1.8	1	7
Home satisfaction	4.5	1.4	1	7
Vehicle satisfaction	4.0	1.6	1	7
Life satisfaction (cont.)	4.3	1.4	1	7
Health satisfaction (cont.)	5.2	1.1	1	7
Income satisfaction (cont.)	3.9	1.7	1	7
Home satisfaction (cont.)	4.6	1.3	1	7
Vehicle satisfaction (cont.)	4.1	1.5	1	7

[illegible][illegible][illegible][illegible]

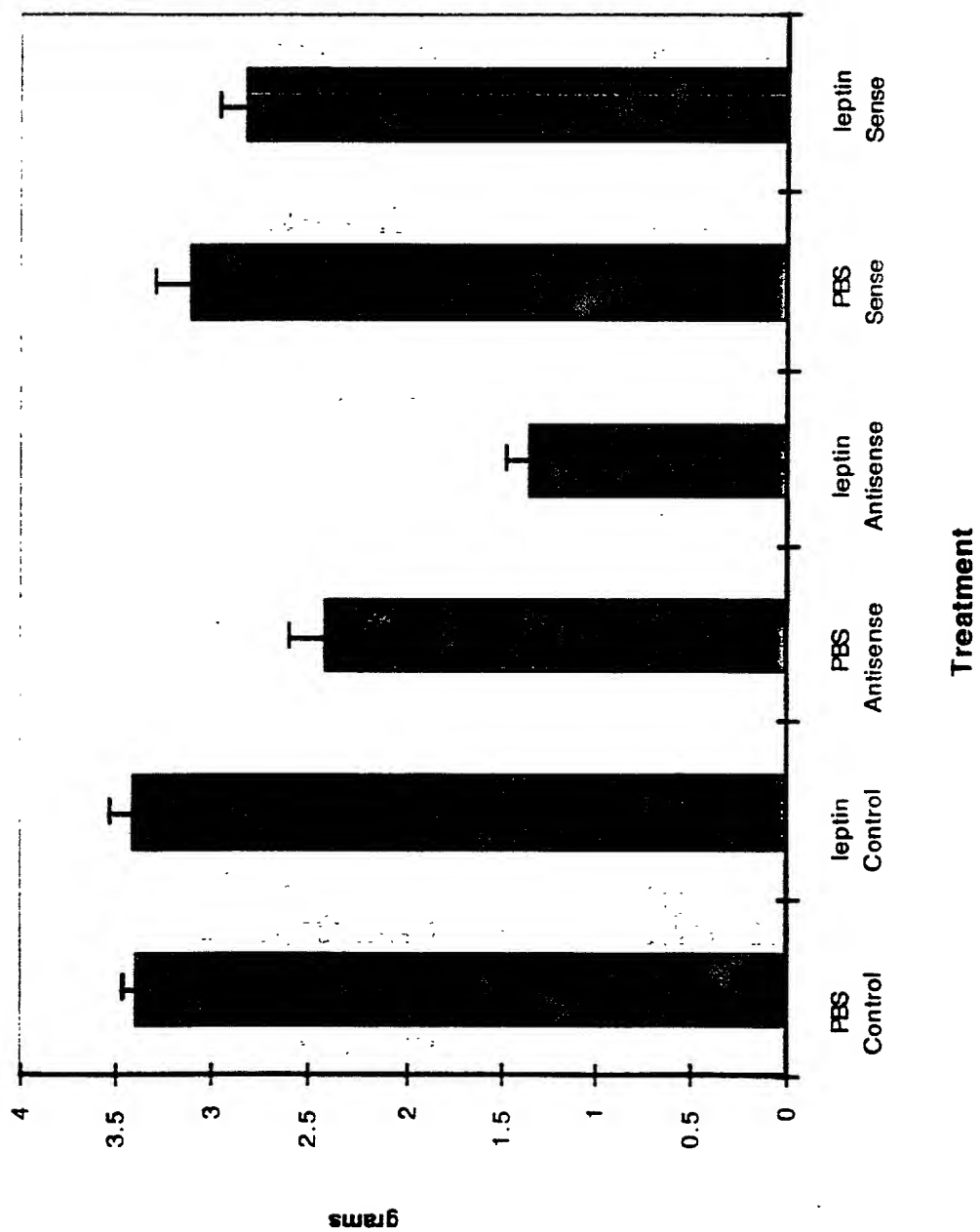


FIG. 11